ENTERED   PCT10

RAW SEQUENCE LISTING   DATE: 04/04/2002
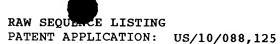PATENT APPLICATION: US/10/088,125   TIME: 15:35:51

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

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   4 <110> APPLICANT: Joelle Thonnard
   6 <120> TITLE OF INVENTION: Novel Compounds
   9 <130> FILE REFERENCE: BM45413
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/088,125
C--> 11 <141> CURRENT FILING DATE: 2002-03-14
  11 <150> PRIOR APPLICATION NUMBER: 99 21692.1
  12 <151> PRIOR FILING DATE: 1999-09-14
  14 <160> NUMBER OF SEQ ID NOS: 10
  16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
  18 <210> SEQ ID NO: 1
  19 <211> LENGTH: 1524
  20 <212> TYPE: DNA
  21 <213> ORGANISM: Moraxella catarrhalis
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  24 atgaaaatct ctacaactgc acgtatcttg acgctctcag cgctggcgat tggcatggca     60
  25 gcttgtagta gcattccaaa gaaaattgac acatcggcac cgattttggc ggtgcctaat    120
  26 gtgcctatga aggatggcta tcaagtctat gatgctgaca ccatcagtgt ggcgcaggca    180
  27 ccaagcgtgg catcactacg ctggcaggaa ttttatacaa atcctaaact tgcagctttg    240
  28 attgagcttg ctttacaaaa taacaaagac ttgcaatctg cggtcttagc cgtgcaatca    300
  29 gcccgtgctc aatatcaaat taccgaagct ggcagcgtgc cacaagtcgg ctcaaatacc    360
  30 agtgtgacac gccaagcgaa taaccgtatc gatgccaatg cttcaaccaa ttatcatgtt    420
  31 gggcttgcga tgagtggtta tgagctggat ttatggggca aagttgccag tcaaaaacag    480
  32 caagcactac atcaatattt ggcaaccaac gccgccaaag acgccgttca gatttcaatc    540
  33 atctcaagcg tcgcccaagg ttacgttaac ttagctcacg ctttggctca aaggcagttg    600
  34 gctgagcaga cgctaaaaac ccgtgaacat gcgatgatga ttacccaaaa gcgttttgaa    660
  35 gcggggattg attctaagtc gccaagtcta caagcagcaa gttcacttga gtcagcacga    720
  36 ttggcagtat atgcagcaga taccagtatc ttaaaagcca aaaatgcgtt acagctgctg    780
  37 attggtcgtc ctgtaccaaa cgagctacta ccagcgatag atgccagtat gcatatgggt    840
  38 catattacca cacagacatt gtttagtgca ggtttgccca gcgagctttt atattatcgc    900
  39 ccagatatta tgcaggctga gcatcgccta aaagcagcag gtgcaaatat caatgtggca    960
  40 cgcgctgctt attttccgtc gattcgttta tcatctaatg tgggatttag tagtaacagt   1020
  41 ttgaataact tatttgaatc aagtgctttg ggctggtctt ttgggcctgc gattagcttg   1080
  42 cctatctttg atgcaggcag tcgccgtgcc aatcatgaga tggcgcaagt tgctcagcag   1140
  43 tcggcattgg tggattatga aaaagctatt caaaatgcct ttaaagaagt gtcggatgtt   1200
  44 ttagctgagc gtgcaacttt aggcttgcgt cttgatgccc agattcgcct tcaggataat   1260
  45 taccgtcaaa cttatgatat cgcttatgca agatttcgtt ctggattgga taattatctg   1320
  46 acggtactgg acgccgagcg gtctttattt attaatcagc aaaatatact acagcttgaa   1380
  47 cttgccaagt tagtcagcca aatccagcta taccaagcat ggggcggcgg tgcaagctta   1440
  48 actgctgagc aaatcacaga atttaatcgt cagcgtgaag ccatgcgtcc agccatgctg   1500
  49 atggatgacg aaaaatccca atag                                         1524
  51 <210> SEQ ID NO: 2
  52 <211> LENGTH: 506
  53 <212> TYPE: PRT
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/088,125

DATE: 04/04/2002
TIME: 15:35:51

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

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56 <400> SEQUENCE: 2
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58 1               5                   10              15
59 Ile Gly Met Ala Ala Cys Ser Ser Ile Pro Lys Lys Ile Asp Thr Ser
60          20              25              30
61 Ala Pro Ile Leu Ala Val Pro Asn Val Pro Met Lys Asp Gly Tyr Gln
62          35              40              45
63 Val Tyr Asp Ala Asp Thr Ile Ser Val Ala Gln Ala Pro Ser Val Ala
64      50              55              60
65 Ser Leu Arg Trp Gln Glu Phe Tyr Thr Asn Pro Lys Leu Ala Ala Leu
66 65              70              75              80
67 Ile Glu Leu Ala Leu Gln Asn Asn Lys Asp Leu Gln Ser Ala Val Leu
68              85              90              95
69 Ala Val Gln Ser Ala Arg Ala Gln Tyr Gln Ile Thr Glu Ala Gly Ser
70              100             105             110
71 Val Pro Gln Val Gly Ser Asn Thr Ser Val Thr Arg Gln Ala Asn Asn
72          115             120             125
73 Arg Ile Asp Ala Asn Ala Ser Thr Asn Tyr His Val Gly Leu Ala Met
74      130             135             140
75 Ser Gly Tyr Glu Leu Asp Leu Trp Gly Lys Val Ala Ser Gln Lys Gln
76 145             150             155             160
77 Gln Ala Leu His Gln Tyr Leu Ala Thr Asn Ala Ala Lys Asp Ala Val
78              165             170             175
79 Gln Ile Ser Ile Ile Ser Ser Val Ala Gln Gly Tyr Val Asn Leu Ala
80              180             185             190
81 His Ala Leu Ala Gln Arg Gln Leu Ala Glu Gln Thr Leu Lys Thr Arg
82              195             200             205
83 Glu His Ala Met Met Ile Thr Gln Lys Arg Phe Ala Gly Ile Asp Ser
84      210             215             220
85 Lys Ser Pro Ser Leu Gln Ala Ala Ser Ser Leu Glu Ser Ala Arg Leu
86 225             230             235             240
87 Ala Val Tyr Ala Ala Asp Thr Ser Ile Leu Lys Ala Lys Asn Ala Leu
88              245             250             255
89 Gln Leu Leu Ile Gly Arg Pro Val Pro Asn Glu Leu Leu Pro Ala Ile
90              260             265             270
91 Asp Ala Ser Met His Met Gly His Ile Thr Thr Gln Thr Leu Phe Ser
92              275             280             285
93 Ala Gly Leu Pro Ser Glu Leu Leu Tyr Tyr Arg Pro Asp Ile Met Gln
94      290             295             300
95 Ala Glu His Arg Leu Lys Ala Ala Gly Ala Asn Ile Asn Val Ala Arg
96 305             310             315             320
97 Ala Ala Tyr Phe Pro Ser Ile Arg Leu Ser Ser Asn Val Gly Phe Ser
98              325             330             335
99 Ser Asn Ser Leu Asn Asn Leu Phe Glu Ser Ser Ala Leu Gly Trp Ser
100             340             345             350
101 Phe Gly Pro Ala Ile Ser Leu Pro Ile Phe Asp Ala Gly Ser Arg Arg
102         355             360             365
103 Ala Asn His Glu Met Ala Gln Val Ala Gln Gln Ser Ala Leu Val Asp
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Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

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104      370                      375                   380
105 Tyr Glu Lys Ala Ile Gln Asn Ala Phe Lys Glu Val Ser Asp Val Leu
106 385                  390                   395                   400
107 Ala Glu Arg Ala Thr Leu Gly Leu Arg Leu Asp Ala Gln Ile Arg Leu
108              405                   410                   415
109 Gln Asp Asn Tyr Arg Gln Thr Tyr Asp Ile Ala Tyr Ala Arg Phe Arg
110              420                   425                   430
111 Ser Gly Leu Asp Asn Tyr Leu Thr Val Leu Asp Ala Glu Arg Ser Leu
112          435                   440                   445
113 Phe Ile Asn Gln Gln Asn Ile Leu Gln Leu Glu Leu Ala Lys Leu Val
114      450                   455                   460
115 Ser Gln Ile Gln Leu Tyr Gln Ala Leu Gly Gly Gly Ala Ser Leu Thr
116 465                  470                   475                   480
117 Ala Glu Gln Ile Thr Glu Phe Asn Arg Gln Arg Glu Ala Met Arg Pro
118              485                   490                   495
119 Ala Met Leu Met Asp Asp Glu Lys Ser Gln
120          500                   505
122 <210> SEQ ID NO: 3
123 <211> LENGTH: 1521
124 <212> TYPE: DNA
125 <213> ORGANISM: Moraxella catarrhalis
127 <400> SEQUENCE: 3
128 atgaaaatct ctacaactgc acgtatcttg acgctctcag cgctggcgat tggcatggca        60
129 gcttgtagta gcattccaaa gaaaattgac acatcggcac cgattttggc ggtgcctaat       120
130 gtgcctatga aggatggcta tcaagtctat gatgctgaca ccatcagtgt ggcgcaggca       180
131 ccaagcgtgg catcactacg ctggcaggaa ttttatacaa atcctaaact tgcagctttg       240
132 attgagcttg ctttacaaaa taacaaagac ttgcaatctg cggtcttagc cgtgcaatca       300
133 gcccgtgctc aatatcaaat taccgaagct ggcagcgtgc cacaagtcgg ctcaaatacc       360
134 agtgtgacac gccaagcgaa taaccgtatc gatgccaatg cttcaaccaa ttatcatgtt       420
135 gggcttgcga tgagtggtta tgagctggat ttatggggca aagttgccag tcaaaaacag       480
136 caagcactac atcaatattt ggcaaccaac gccgccaaag acgccgttca gatttcaatc       540
137 atctcaagcg tcgcccaagg ttacgttaac ttagctcacg ctttggctca aaggcagttg       600
138 gctgagcaga cgctaaaaac ccgtgaacat gcgatgatga ttacccaaaa gcgttttgaa       660
139 gcggggattg attctaagtc gccaagtcta caagcagcaa gttcacttga gtcagcacga       720
140 ttggcagtat atgcagcaga taccagtatc ttaaaagcca aaaatgcgtt acagctgctg       780
141 attggtcgtc ctgtaccaaa cgagctacta ccagcgatag atgccagtat gcatatgggt       840
142 catattacca cacagacatt gtttagtgca ggtttgccca gcgagctttt atattatcgc       900
143 ccagatatta tgcaggctga gcatcgccta aaagcagcag gtgcaaatat caatgtggca       960
144 cgcgctgctt attttccgtc gattcgttta tcatctaatg tgggatttag tagtaacagt      1020
145 ttgaataact tatttgaatc aagtgctttg ggctggtctt ttgggcctgc gattagcttg      1080
146 cctatctttg atgcaggcag tcgccgtgcc aatcatgaga tggcgcaagt tgctcagcag      1140
147 tcggcattgg tggattatga aaaagctatt caaaatgcct ttaaagaagt gtcggatgtt      1200
148 ttagctgagc gtgcaacttt aggcttgcgt cttgatgccc agattcgcct tcaggataat      1260
149 taccgtcaaa cttatgatat cgcttatgca agatttcgtt ctggattgga taattatctg      1320
150 acggtactgg acgccgagcg gtctttattt attaatcagc aaaatatact acagcttgaa      1380
151 cttgccaagt tagtcagcca aatccagcta taccaagcat tgggcggcgg tgcaagctta      1440
152 actgctgagc aaatcacaga atttaatcgt cagcgtgaag ccatgcgtcc agccatgctg      1500
153 atggatgacg aaaaatccca a                                                1521
155 <210> SEQ ID NO: 4
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/088,125

DATE: 04/04/2002
TIME: 15:35:51

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

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156 <211> LENGTH: 506
157 <212> TYPE: PRT
158 <213> ORGANISM: Moraxella catarrhalis
160 <400> SEQUENCE: 4
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163 Ile Gly Met Ala Ala Cys Ser Ser Ile Pro Lys Lys Ile Asp Thr Ser
164             20                  25                  30
165 Ala Pro Ile Leu Ala Val Pro Asn Val Pro Met Lys Asp Gly Tyr Gln
166             35                  40                  45
167 Val Tyr Asp Ala Asp Thr Ile Ser Val Ala Gln Ala Pro Ser Val Ala
168     50                  55                  60
169 Ser Leu Arg Trp Gln Glu Phe Tyr Thr Asn Pro Lys Leu Ala Ala Leu
170 65                  70                  75                  80
171 Ile Glu Leu Ala Leu Gln Asn Asn Lys Asp Leu Gln Ser Ala Val Leu
172             85                  90                  95
173 Ala Val Gln Ser Ala Arg Ala Gln Tyr Gln Ile Thr Glu Ala Gly Ser
174             100                 105                 110
175 Val Pro Gln Val Gly Ser Asn Thr Ser Val Thr Arg Gln Ala Asn Asn
176             115                 120                 125
177 Arg Ile Asp Ala Asn Ala Ser Thr Asn Tyr His Val Gly Leu Ala Met
178         130                 135                 140
179 Ser Gly Tyr Glu Leu Asp Leu Trp Gly Lys Val Ala Ser Gln Lys Gln
180 145                 150                 155                 160
181 Gln Ala Leu His Gln Tyr Leu Ala Thr Asn Ala Ala Lys Asp Ala Val
182             165                 170                 175
183 Gln Ile Ser Ile Ile Ser Ser Val Ala Gln Gly Tyr Val Asn Leu Ala
184             180                 185                 190
185 His Ala Leu Ala Gln Arg Gln Leu Ala Glu Gln Thr Leu Lys Thr Arg
186         195                 200                 205
187 Glu His Ala Met Met Ile Thr Gln Lys Arg Phe Ala Gly Ile Asp Ser
188     210                 215                 220
189 Lys Ser Pro Ser Leu Gln Ala Ala Ser Ser Leu Glu Ser Ala Arg Leu
190 225                 230                 235                 240
191 Ala Val Tyr Ala Ala Asp Thr Ser Ile Leu Lys Ala Lys Asn Ala Leu
192             245                 250                 255
193 Gln Leu Leu Ile Gly Arg Pro Val Pro Asn Glu Leu Leu Pro Ala Ile
194             260                 265                 270
195 Asp Ala Ser Met His Met Gly His Ile Thr Thr Gln Thr Leu Phe Ser
196         275                 280                 285
197 Ala Gly Leu Pro Ser Glu Leu Leu Tyr Tyr Arg Pro Asp Ile Met Gln
198     290                 295                 300
199 Ala Glu His Arg Leu Lys Ala Ala Gly Ala Asn Ile Asn Val Ala Arg
200 305                 310                 315                 320
201 Ala Ala Tyr Phe Pro Ser Ile Arg Leu Ser Ser Asn Val Gly Phe Ser
202             325                 330                 335
203 Ser Asn Ser Leu Asn Asn Leu Phe Glu Ser Ser Ala Leu Gly Trp Ser
204         340                 345                 350
205 Phe Gly Pro Ala Ile Ser Leu Pro Ile Phe Asp Ala Gly Ser Arg Arg
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/088,125
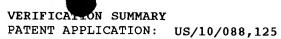
DATE: 04/04/2002
TIME: 15:35:51

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

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206          355                    360                    365
207 Ala Asn His Glu Met Ala Gln Val Ala Gln Gln Ser Ala Leu Val Asp
208     370                375                380
209 Tyr Glu Lys Ala Ile Gln Asn Ala Phe Lys Glu Val Ser Asp Val Leu
210 385                390                395                400
211 Ala Glu Arg Ala Thr Leu Gly Leu Arg Leu Asp Ala Gln Ile Arg Leu
212             405                410                415
213 Gln Asp Asn Tyr Arg Gln Thr Tyr Asp Ile Ala Tyr Ala Arg Phe Arg
214             420                425                430
215 Ser Gly Leu Asp Asn Tyr Leu Thr Val Leu Asp Ala Glu Arg Ser Leu
216             435                440                445
217 Phe Ile Asn Gln Gln Asn Ile Leu Gln Leu Glu Leu Ala Lys Leu Val
218     450                455                460
219 Ser Gln Ile Gln Leu Tyr Gln Ala Leu Gly Gly Gly Ala Ser Leu Thr
220 465                470                475                480
221 Ala Glu Gln Ile Thr Glu Phe Asn Arg Gln Arg Glu Ala Met Arg Pro
222             485                490                495
223 Ala Met Leu Met Asp Asp Glu Lys Ser Gln
224             500                505
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227 <211> LENGTH: 21
228 <212> TYPE: DNA
229 <213> ORGANISM: Artificial Sequence
231 <220> FEATURE:
232 <223> OTHER INFORMATION: primer
234 <400> SEQUENCE: 5
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237 <210> SEQ ID NO: 6
238 <211> LENGTH: 22
239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial Sequence
242 <220> FEATURE:
243 <223> OTHER INFORMATION: primer
245 <400> SEQUENCE: 6
246 tggtcgtcct gtaccaaacg ag                                      22
248 <210> SEQ ID NO: 7
249 <211> LENGTH: 17
250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: primer
256 <400> SEQUENCE: 7
257 gtaaaacgac ggccagt                                            17
259 <210> SEQ ID NO: 8
260 <211> LENGTH: 17
261 <212> TYPE: DNA
262 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: primer
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VERIFICATION SUMMARY   DATE: 04/04/2002
PATENT APPLICATION: US/10/088,125   TIME: 15:35:52

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04042002\J088125.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date